## **RAW SEQUENCE LISTING**

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number:	10/552,291
Source:	PUTIO
Date Processed by STIC:	10/18/05

## ENTERED



PCT

RAW SEQUENCE LISTING DATE: 10/18/2005 PATENT APPLICATION: US/10/552,291 TIME: 10:11:21

Input Set: A:\05-10-03 Sequence Listing.txt
Output Set: N:\CRF4\10182005\J552291.raw

```
3 <110> APPLICANT: REGEN Biotech, Inc.
     5 <120> TITLE OF INVENTION: Use of a peptide that interacts with alpha v beta3 integrin
of
              endothelial cell
      8 <130> FILE REFERENCE: OP04-1024
C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/552,291
C--> 10 <141> CURRENT FILING DATE: 2005-10-03
     10 <150> PRIOR APPLICATION NUMBER: KR 10-2003-0021065
     11 <151> PRIOR FILING DATE: 2003-04-03
     13 <160> NUMBER OF SEQ ID NOS: 27
     15 <170> SOFTWARE: KopatentIn 1.71
     17 <210> SEQ ID NO: 1
     18 <211> LENGTH: 683
    19 <212> TYPE: PRT
    20 <213> ORGANISM: Homo sapiens
     22 <400> SEQUENCE: 1
    23 Met Ala Leu Phe Val Arg Leu Leu Ala Leu Ala Leu Ala Leu
                          5
    24 1
                                             10
     26 Gly Pro Ala Ala Thr Leu Ala Gly Pro Ala Lys Ser Pro Tyr Gln Leu
                    20
    29 Val Leu Gln His Ser Arg Leu Arg Gly Arg Gln His Gly Pro Asn Val
                                     40
     32 Cys Ala Val Gln Lys Val Ile Gly Thr Asn Arg Lys Tyr Phe Thr Asn
    35 Cys Lys Gln Trp Tyr Gln Arg Lys Ile Cys Gly Lys Ser Thr Val Ile
                            70
     38 Ser Tyr Glu Cys Cys Pro Gly Tyr Glu Lys Val Pro Gly Glu Lys Gly
    41 Cys Pro Ala Ala Leu Pro Leu Ser Asn Leu Tyr Glu Thr Leu Gly Val
                                        105
                    100
     44 Val Gly Ser Thr Thr Thr Gln Leu Tyr Thr Asp Arg Thr Glu Lys Leu
                                    120
     45
    47 Arg Pro Glu Met Glu Gly Pro Gly Ser Phe Thr Ile Phe Ala Pro Ser
    50 Asn Glu Ala Trp Ala Ser Leu Pro Ala Glu Val Leu Asp Ser Leu Val
                            150
                                                155
    53 Ser Asn Val Asn Ile Glu Leu Leu Asn Ala Leu Arg Tyr His Met Val
                                            170
    56 Gly Arg Arg Val Leu Thr Asp Glu Leu Lys His Gly Met Thr Leu Thr
                    180
                                        185
                                                            190
    59 Ser Met Tyr Gln Asn Ser Asn Ile Gln Ile His His Tyr Pro Asn Gly
               195
                                    200
                                                        205
    62 Ile Val Thr Val Asn Cys Ala Arg Leu Leu Lys Ala Asp His His Ala
```

220

215

Input Set : A:\05-10-03 Sequence Listing.txt
Output Set: N:\CRF4\10182005\J552291.raw

65 Thr Asn Gly Val Val His Leu Ile Asp Lys Val Ile Ser Thr Ile Thr 66 225 68 Asn Asn Ile Gln Gln Ile Ile Glu Ile Glu Asp Thr Phe Glu Thr Leu 245 250 71 Arg Ala Ala Val Ala Ala Ser Gly Leu Asn Thr Met Leu Glu Gly Asn 265 74 Gly Gln Tyr Thr Leu Leu Ala Pro Thr Asn Glu Ala Phe Glu Lys Ile 275 280 77 Pro Ser Glu Thr Leu Asn Arg Ile Leu Gly Asp Pro Glu Ala Leu Arg 295 300 80 Asp Leu Leu Asn Asn His Ile Leu Lys Ser Ala Met Cys Ala Glu Ala 310 315 83 Ile Val Ala Gly Leu Ser Val Glu Thr Leu Glu Gly Thr Thr Leu Glu 325 330 86 Val Gly Cys Ser Gly Asp Met Leu Thr Ile Asn Gly Lys Ala Ile Ile 89 Ser Asn Lys Asp Ile Leu Ala Thr Asn Gly Val Ile His Tyr Ile Asp 355 360 92 Glu Leu Leu Ile Pro Asp Ser Ala Lys Thr Leu Phe Glu Leu Ala Ala 375 95 Glu Ser Asp Val Ser Thr Ala Ile Asp Leu Phe Arg Gln Ala Gly Leu 395 390 98 Gly Asn His Leu Ser Gly Ser Glu Arg Leu Thr Leu Leu Ala Pro Leu 405 410 101 Asn Ser Val Phe Lys Asp Gly Thr Pro Pro Ile Asp Ala His Thr Arg 425 420 104 Asn Leu Leu Arg Asn His Ile Ile Lys Asp Gln Leu Ala Ser Lys Tyr 435 107 Leu Tyr His Gly Gln Thr Leu Glu Thr Leu Gly Gly Lys Lys Leu Arg 455 110 Val Phe Val Tyr Arg Asn Ser Leu Cys Ile Glu Asn Ser Cys Ile Ala 470 475 111 465 113 Ala His Asp Lys Arg Gly Arg Tyr Gly Thr Leu Phe Thr Met Asp Arg 485 490 116 Val Leu Thr Pro Pro Met Gly Thr Val Met Asp Val Leu Lys Gly Asp 117 500 505 119 Asn Arg Phe Ser Met Leu Val Ala Ala Ile Gln Ser Ala Gly Leu Thr 520 122 Glu Thr Leu Asn Arg Glu Gly Val Tyr Thr Val Phe Ala Pro Thr Asn 535 540 125 Glu Ala Phe Arg Ala Leu Pro Pro Arg Glu Arg Ser Arg Leu Leu Gly 550 555 128 Asp Ala Lys Glu Leu Ala Asn Ile Leu Lys Tyr His Ile Gly Asp Glu 570 565 131 Ile Leu Val Ser Gly Gly Ile Gly Ala Leu Val Arg Leu Lys Ser Leu 132 585 134 Gln Gly Asp Lys Leu Glu Val Ser Leu Lys Asn Asn Val Val Ser Val 595 600 137 Asn Lys Glu Pro Val Ala Glu Pro Asp Ile Met Ala Thr Asn Gly Val

Input Set : A:\05-10-03 Sequence Listing.txt
Output Set: N:\CRF4\10182005\J552291.raw

138	_	610	_	_	_		615		_			620				
		His	Val	Ile	Thr	Asn	Val	Leu	Gln	Pro		Ala	Asn	Arg	Pro	
	625					630					635					640
143	Glu	Arg	Gly	Asp	Glu	Leu	Ala	Asp	Ser	Ala	Leu	Glu	Ile	Phe	Lys	Gln
144					645					650					655	
146	Ala	Ser	Ala	Phe	Ser	Arg	Ala	Ser	Gln	Arg	Ser	Val	Arg	Leu	Ala	Pro
147				660					665					670		
149	Val	Tyr	Gln	Lys	Leu	Leu	Glu	Arg	Met	Lys	His					
150			675					680								
153	<21	0> SI	EQ II	ONO:	: 2											
154	54 <211> LENGTH: 103															
155	5 <212> TYPE: PRT															
156	<21	3 > OI	RGAN:	ISM:	Homo	sar	oiens	3								
	<40					1										
						Thr	Ile	Phe	Ala	Pro	Ser	Asn	Glu	Ala	Tro	Ala
160	1		1		5					10					15	
		Leu	Pro	Ala	_	Val	Len	Asp	Ser		Val	Ser	Asn	Val		Tle
163	-			20			<b></b>	1.05	25					30		
	Glu	T.e11	T.e.ii		Δla	Leu	Ara	Tvr		Met	Val	Glv	Ara		Val	Len
166	014	u	35	1.011			••••	40			· u =		45	5	***	200
	Thr	Aen		T.e.11	Lve	His	Glv		Thr	T.e.11	Thr	Ser		Tur	Gln	Δen
169	1111	50	GIU	пси	цуз	1113	55	MCC	****	пси	1111	60	rice	-1-	0111	11011
	Sar		т1Д	Gln	Tla	His		Тиг	Dro	Aen	Glv		Val	Thr	Val	Δen
172		Noii	116	GIII	116	70	1113	T Y L	rio	AŞII	75	116	vai	1111	Val	80
		λla	7 ra	Lou	Lou	Lys	λla	Nan	uic	uic		Thr	λan	Clv	17-1	
175	Cys	AIG	Ar 9	пси	85	цуз	AIG	тар	1115	90	пια	1111	ASII	Gry	95	Vai
	uic	T.011	Tla	λan		17a l	בוד			90					93	
	птр	пеп	116	100	БХР	Val	116							•		•
178	-21	n	20 TI													
	<21															
	<21				31											
	<21				TT			_								
						sa <u>r</u>	)iens	j .								
	<40					*1.	<b>a</b> 1	<b>-1</b>	<b>~1</b>	3	m\	D1	<b>~1</b>	ml	T	7
		TTE	GIII	GIII		Ile	GIU	тте	GIU		THE	Pne	GIU	THE		Arg
188	1	<b>31</b> -	777	77-	5	0	<b>a</b> 1	T	7	10	Ma.L	<b>+</b>	a1	<b>a</b> 1	15	<b>C1</b>
	Ата	Ala	vai		АТА	Ser	GIY	ьeu		Thr	Met	ьeu	GIU		ASII	GIY
191	<b>~</b> 1-	m	m\	20	<b>T</b>	<b>71</b> -	D	m1	25	<b>a</b> 1	<b>77</b> -	D1	~1	30	<b>7</b> 7.	D
	GIn	ıyr		ьeu	Leu	Ala	Pro		Asn	GIU	Ата	Pne		гаг	ше	Pro
194	_		35	_	_	_		40		_	_		45	_	_	_
	Ser		Thr	Leu	Asn	Arg		Leu	GIY	Asp	Pro		Ala	Leu	Arg	Asp
197		50			_	_	55			_		60	_		_	
		Leu	Asn	Asn	His	Ile	Leu	Lys	Ser	Ala		Cys	Ala	Glu	Ala	
200						70				_	75		_			80_
	Val	Ala	Gly	Leu		Val	Glu	Thr	Leu		Gly	Thr	Thr	Leu	Glu	Val
203					85					90					95	
205	Gly	Cys	Ser	Gly	Asp	Met	Leu	Thr		Asn	Gly	Lys	Ala	Ile	Ile	Ser
206				100					105					110		
208	Asn	Lys	Asp	Ile	Leu	Ala	Thr	Asn	Gly	Val	Ile	His	Tyr	Ile	Asp	Glu
209			115					120					125			

Input Set : A:\05-10-03 Sequence Listing.txt
Output Set: N:\CRF4\10182005\J552291.raw

```
211 Leu Leu Ile
212
        130
215 <210> SEQ ID NO: 4
216 <211> LENGTH: 129
217 <212> TYPE: PRT
218 <213> ORGANISM: Homo sapiens
220 <400> SEQUENCE: 4
221 Pro Asp Ser Ala Lys Thr Leu Phe Glu Leu Ala Ala Glu Ser Asp Val
224 Ser Thr Ala Ile Asp Leu Phe Arg Gln Ala Gly Leu Gly Asn His Leu
                 20
                                     25
227 Ser Gly Ser Glu Arg Leu Thr Leu Leu Ala Pro Leu Asn Ser Val Phe
                                 40
230 Lys Asp Gly Thr Pro Pro Ile Asp Ala His Thr Arg Asn Leu Leu Arg
233 Asn His Ile Ile Lys Asp Gln Leu Ala Ser Lys Tyr Leu Tyr His Gly
236 Gln Thr Leu Glu Thr Leu Gly Gly Lys Lys Leu Arg Val Phe Val Tyr
237
239 Arg Asn Ser Leu Cys Ile Glu Asn Ser Cys Ile Ala Ala His Asp Lys
                                    105
242 Arg Gly Arg Tyr Gly Thr Leu Phe Thr Met Asp Arg Val Leu Thr Pro
243
            115
                                120
                                                     125
245 Pro
249 <210> SEQ ID NO: 5
250 <211> LENGTH: 131
251 <212> TYPE: PRT
252 <213> ORGANISM: Homo sapiens
254 <400> SEQUENCE: 5
255 Met Gly Thr Val Met Asp Val Leu Lys Gly Asp Asn Arg Phe Ser Met
256 1
258 Leu Val Ala Ala Ile Gln Ser Ala Gly Leu Thr Glu Thr Leu Asn Arg
                 20
261 Glu Gly Val Tyr Thr Val Phe Ala Pro Thr Asn Glu Ala Phe Arg Ala
            35
                                 40
264 Leu Pro Pro Arg Glu Arg Ser Arg Leu Leu Gly Asp Ala Lys Glu Leu
                             55
267 Ala Asn Ile Leu Lys Tyr His Ile Gly Asp Glu Ile Leu Val Ser Gly
270 Gly Ile Gly Ala Leu Val Arg Leu Lys Ser Leu Gln Gly Asp Lys Leu
                     85
273 Glu Val Ser Leu Lys Asn Asn Val Val Ser Val Asn Lys Glu Pro Val
                                    105
276 Ala Glu Pro Asp Ile Met Ala Thr Asn Gly Val Val His Val Ile Thr
                                120
277
            115
279 Asn Val Leu
280
        130
283 <210> SEQ ID NO: 6
284 <211> LENGTH: 85
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Input Set : A:\05-10-03 Sequence Listing.txt
Output Set: N:\CRF4\10182005\J552291.raw

```
285 <212> TYPE: PRT
286 <213> ORGANISM: Homo sapiens
288 <400> SEQUENCE: 6
289 Arg Ala Leu Pro Pro Arg Glu Arg Ser Arg Leu Leu Gly Asp Ala Lys
292 Glu Leu Ala Asn Ile Leu Lys Tyr His Ile Gly Asp Glu Ile Leu Val
               20
                                     25
295 Ser Gly Gly Ile Gly Ala Leu Val Arg Leu Lys Ser Leu Gln Gly Asp
           35
                                 40
298 Lys Leu Glu Val Ser Leu Lys Asn Asn Val Val Ser Val Asn Lys Glu
      50
                            55
301 Pro Val Ala Glu Pro Asp Ile Met Ala Thr Asn Gly Val Val His Val
304 Ile Thr Asn Val Leu
305
308 <210> SEQ ID NO: 7
309 <211> LENGTH: 119
310 <212> TYPE: PRT
311 <213> ORGANISM: Homo sapiens
313 <400> SEQUENCE: 7
314 Met Gly Thr Val Met Asp Val Leu Lys Gly Asp Asn Arg Phe Ser Met
317 Leu Val Ala Ala Ile Gln Ser Ala Gly Leu Thr Glu Thr Leu Asn Arg
                20
                                     25
320 Glu Gly Val Tyr Thr Val Phe Ala Pro Thr Asn Glu Ala Phe Arg Ala
            35
                                 40
323 Leu Pro Pro Arg Glu Arg Ser Arg Leu Leu Gly Asp Ala Lys Glu Leu
326 Ala Asn Ile Leu Lys Tyr His Ile Gly Asp Glu Ile Leu Val Ser Gly
329 Gly Ile Gly Ala Leu Val Arg Leu Lys Ser Leu Gln Gly Asp Lys Leu
                    85
                                        90
332 Glu Val Ser Leu Lys Asn Asn Val Val Ser Val Asn Lys Glu Pro Val
               100
                                   105
335 Ala Glu Pro Asp Ile Met Ala
336
           115
339 <210> SEQ ID NO: 8
340 <211> LENGTH: 113
341 <212> TYPE: PRT
342 <213 > ORGANISM: Homo sapiens
344 <400> SEQUENCE: 8
345 Met Gly Thr Val Met Asp Val Leu Lys Gly Asp Asn Arg Phe Ser Met
346 1 5
348 Leu Val Ala Ala Ile Gln Ser Ala Gly Leu Thr Glu Thr Leu Asn Arg
                                     25
351 Glu Gly Val Tyr Thr Val Phe Ala Pro Thr Asn Glu Ala Phe Arg Ala
                                 40
354 Leu Pro Pro Arg Glu Arg Ser Arg Leu Leu Gly Asp Ala Lys Glu Leu
                            55
```

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/552,291

DATE: 10/18/2005 TIME: 10:11:22

Input Set : A:\05-10-03 Sequence Listing.txt
Output Set: N:\CRF4\10182005\J552291.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application No

L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date